

FIGURE 1

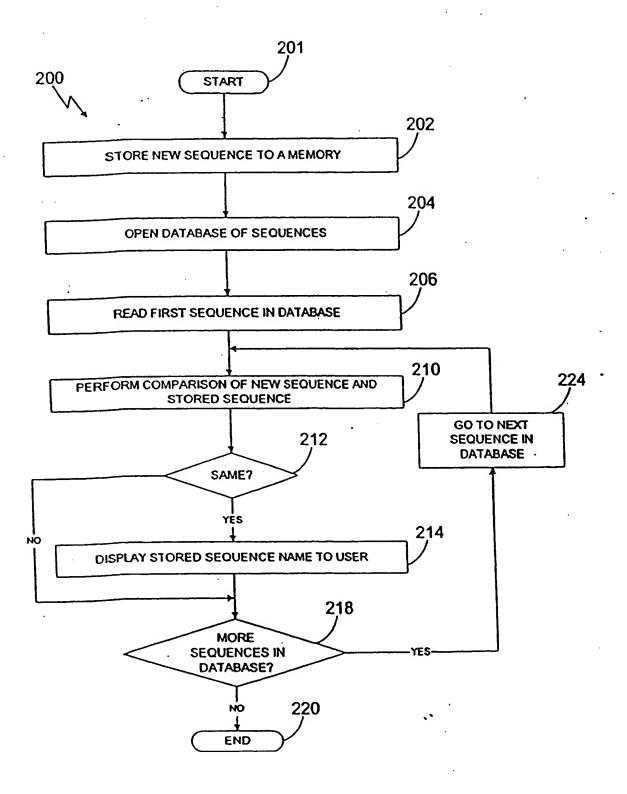
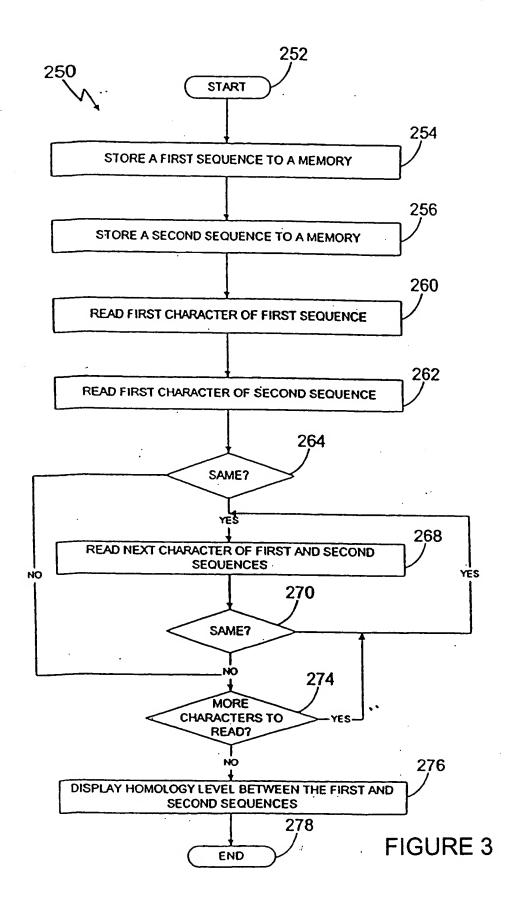
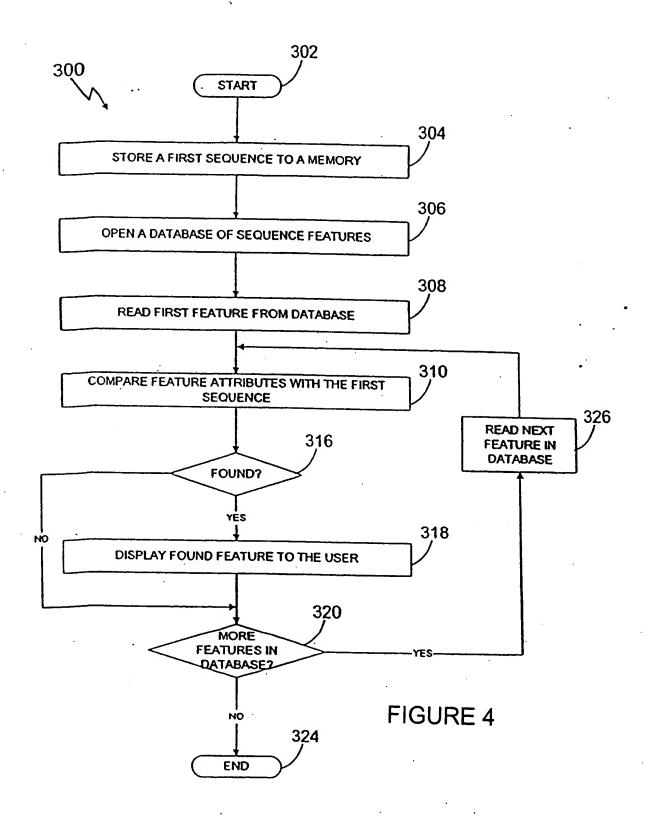


FIGURE 2





Properties of Diversa Fluorescent Proteins

DVSACyan

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28.6		57.3	487	507	0.61	98,200
	2	D.	4	4)	0	86
	25.9	51.8	448(463)	491	0.76	18,900
acids	Calculated subunit mass (kDa)	Total mass (kDa)	Excitation maximum (nm)	Emission maximum (nm)	Quantum yield	Extinction coefficient (M ⁻¹ cm ⁻¹)

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Excitation Maxima

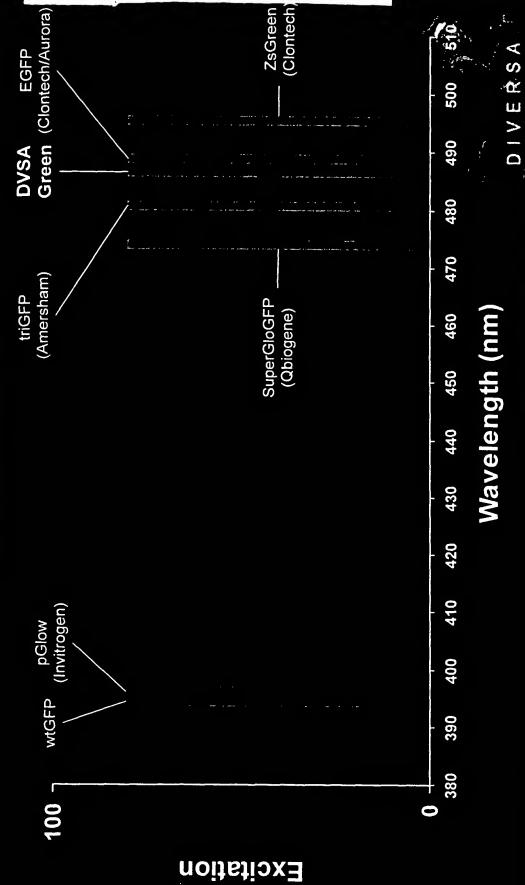
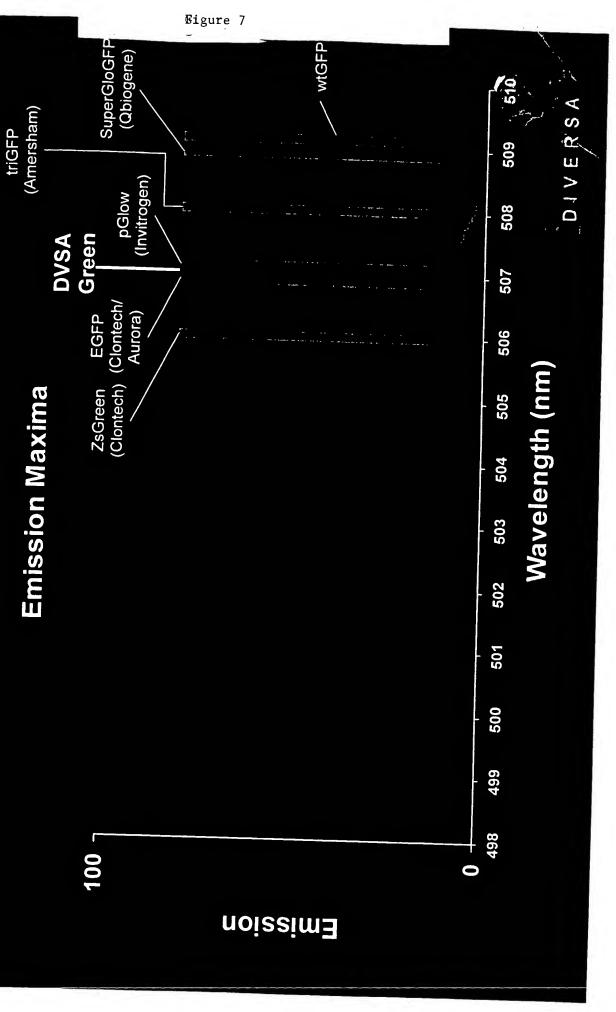


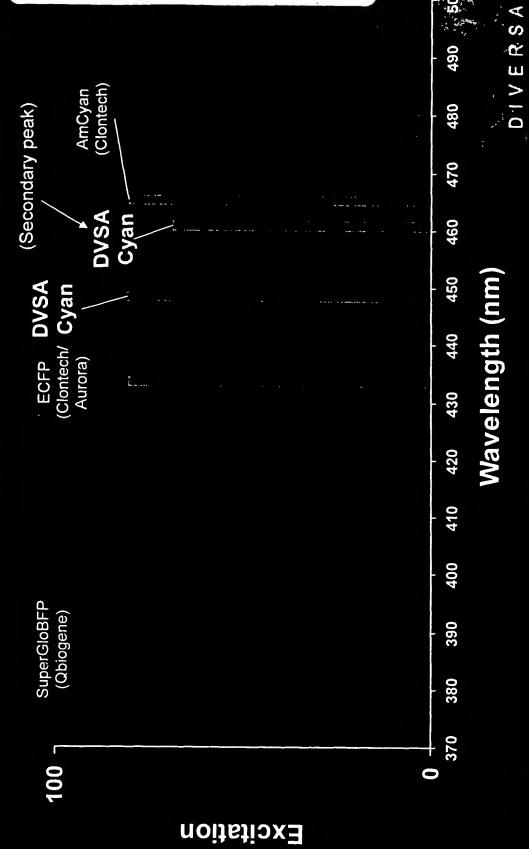
Figure 6

DVSAGreen vs. Other GFPs



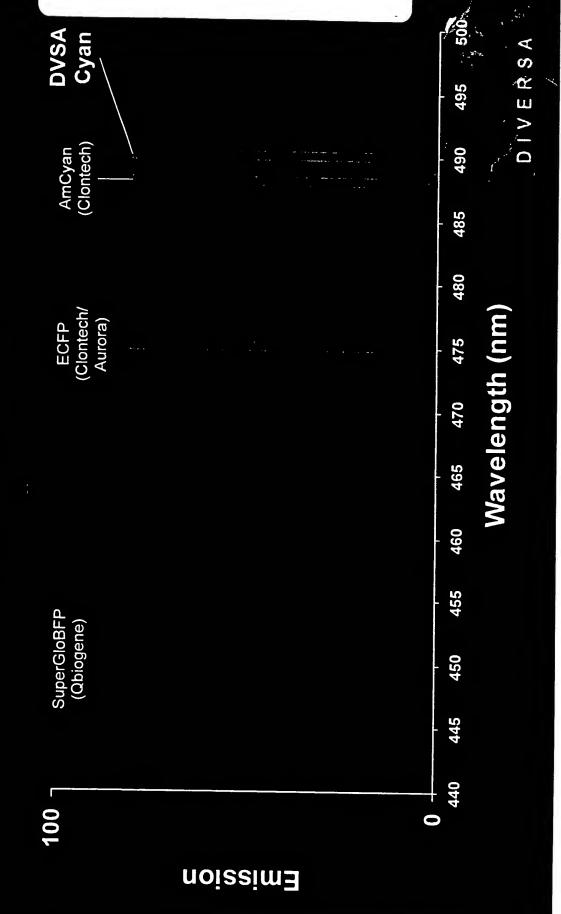
DVSACyan vs. Other Blue/Cyan FPs

Excitation Maxima

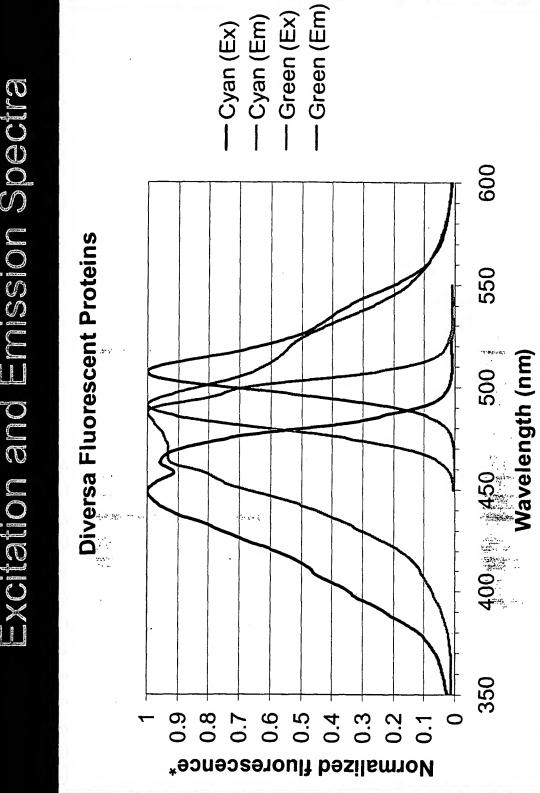


DVSACyan vs. Other Blue/Cyan FPs

Emission Maxima



Excitation and Emission Spectra



Spectra normalized to the peak excitation and emission fluorescence for each protein

DVSAGreen protein is brighter than EGFP

Quantum yield

Extinction coefficient Relative brightness* (M-1 cm-1)

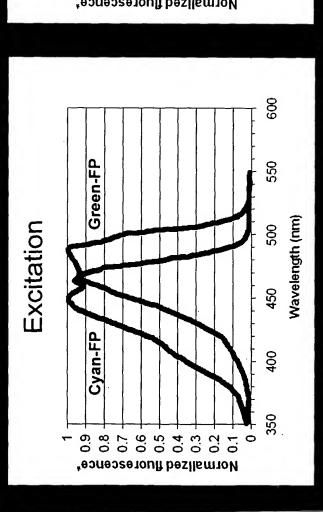
wtGFP	0.77-0.80 1,2	21,600-27,600 1.2	
EGFP	0.6-0.7 3,4	39,200-55,900 ^{3,4}	1.42-1.77
DVS/Groon			2.5-3.5
AmCyan	0.24 ²	40,000 ²	0.43-0.58
DVSACyan	0.76	18,900	0.65-0.88

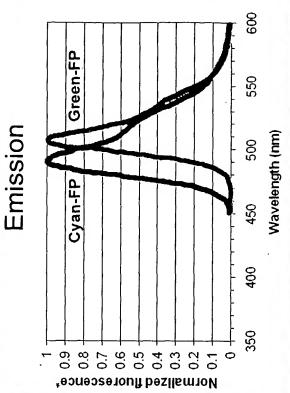
^{*} Relative brightness (maximal extinction coefficient multiplied by quantum yield) as compared to wtGFP

¹Taken from Heim and Tsien, Current Biology 1996 ²Taken from Matz et al, Nature Biotechnology, 1999 ³Taken from Zimmer, Chemical Reviews, 2002 ⁴Taken from Remington, Nature Biotechnology, 2002

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Excitation and Emission Spectra





*Spectra normalized to the peak excitation and emission fluorescence for each protein

Figure 12



Relative Brightness

	Quantum Yield	Extinction Coefficient (M ⁻¹ cm ⁻¹)	Relative Brightne
biscovery Porar "Crean PP	0,61	100 Sept.	2,7-3,6
Wild type AvGFP	0.77-0.80 1.2	21,600-27,600 13	1
EGFP	0.6-0.7	39,200-55,900 **	1.42-1.77
pGlow	0.79	30,000 3	1.1-1.4
Discovery Point Conn. IP	6,7,6	IS, MAN	0865-0833
AmCyan	0.24 ²	40,000 ²	0.43-0.58
ECFP	0.4 ³	32,500	0.59-0.78

Relative brightness (maximal extinction coeff cient multiplied by quantum yield) as compared to wtAvGFP, +Measured per chromophore

Heim and Tsien, Current Biology 1996
Matz et al. Nature Biotechnology, 1999
Remington, N

3. Zimmer, Chemical Reviews, 20024. Remington, Nature Biotechnology, 2002

Figure 13



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Summary of Diversa's DiscoveryPoint™ Fluorescent Proteins
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Excitation/Emission max (nm)	487/507	448(463)/491
Stoke's shift (nm)	20	43(28)
Maturation time	Within 1 hour	Within 1 hour
Quantum yield	0.61	0.76
Extinction coefficient (M ⁻¹ cm ⁻¹)	98,200	18,900
Thermostable to 80°C	Yes	Yes
, # of amino acids	228	227
Calculated subunit mass (kDa)	26.0	25.9
Total mass (kDa) - dimers	52.0	51.8

Figure 14

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NO: 31 ide) of	segment (start-	Overhangs on start-stop	start-CATA/GTAT	CATA/GTAT-TCCT/AGGA	TCCT/AGGA-GGA/CCT	GGA/CCT-TTI/AAA	TTT/AAA-AGG/TCC	AGG/TCC-CTC/GAG	CTC/GAG-ACCA/TGGT	ACCA/TGGT-CCC/GGG	CCC/GGG-CT/GA	CT/GA-AAG/TTC		TTC/AAG-CCT/GGA	CCT/GGA-CATC/GTAG		GGA/CCT-GG/CC	GG/CC-AAG/TTC	AAG/TTC-GA/CT	
SEQ ID NO: 31 Nucleotide	location of	segmen	stop)	143	48-92	97-142	146-167	171-205	209-246	250-274	279-313	317-367	370-415	419-452	456-491	495-534	539-584	588-617	620-659	663-710	743 7EE
			Overhangs on start-stop stop	start-GGA/CCT			GGA/CCT-TTT/AAA	TTT/AAA-AGG/TCC	AGG/TCC-CTC/GAG	CTC/GAG-ACCA/TGGT	ACCA/TGGT-CCC/GGG	CCC/GGG-CT/GA	CT/GA-AAG/TTC	AAG/TTC-TTC/AAG	TTC/AAG-CCT/GGA	CCT/GGA-CATC/GTAG	CATC/GTAG-GGA/CCT	GGA/CCT-GG/CC	GG/CC-AAG/TTC	AAG/TTC-GA/CT	CECIONO ECINO
SEQ ID NO: 29 Nucleotide	location of	segment (start-	stop)	1-41			45-66	70-104	108-145	149-173	178-212	216-266	269-314	318-342	346-378	382-421	426-471	475-504	507-546	250-597	000
			Overhangs on start-stop	start-GGA/CCT			GGA/CCT-TTT/AAA	TTT/AAA-AGG/TCC	AGG/TCC-CTC/GAG	CTC/GAG-ACCA/TGGT	ACCA/TGGT-CCC/GGG	CCC/GGG-CT/GA	CT/GA-AAG/TTC	AAG/TTC-TTC/AAG	TTC/AAG-CCT/GGA	CCT/GGA-CATC/GTAG	CATC/GTAG-GGA/CCT	GGA/CCT-GG/CC	GG/CC-AAG/TTC	AAG/TTC-GA/CT	OFOIO FOI VO
SEQ ID NO: 27 Nucleotide	location of	segm nt (start-	stop) (1-53			27-78	82-116	120-157		190-224	228-275	278-323	327-354	358-393	397-436	441-477	481-500	503-542	546-593	505 630

Figure 15